

## **IN THE FIGURES:**

(A marked-up version of the amendments to the Figures, with changes indicated in RED underlining is enclosed. The amended Formal Drawings are submitted as the clean copy incorporating the amendments indicated below.)

In Figure 1A, please replace the caption with the following caption:

**The inversion sequence of the apo-dystrophin-4 cDNA plus a 10 base-pair region 5' to the start of the inversion sequence (SEQ ID NO: 1A, included in SEQ ID NO: 2).**

In Figure 1B, please replace the caption with the following caption:

**The inversion sequence of the apo-dystrophin-4 cDNA plus the upstream 150 bp from the start of the inversion at 860 to the Hpa I enzyme site (SEQ ID NO: 1B, included in SEQ ID NO: 2)**

In Figure 6, please replace the caption with the following caption:

**FIGURE 6 (SEQ ID NO: 51)**

In Figure 11, please replace lines 1-4 with the following:

|                         |            |            |                                  |
|-------------------------|------------|------------|----------------------------------|
|                         | 1          |            | 50                               |
| (SEQ ID NO 54) Mgen1073 | .....      | .....      | .....                            |
| (SEQ ID NO 55) Hapo1234 | ctagtttcct | attcaatgta | tagtgcacca aaggtcaatt caagagttta |
| Consensus               | -----      | -----      | -----                            |

In Figure 12B, please replace the Figure with the following:

-70 bp from 5' end of apo-4

|

Inr = GCCC TCAT TCTG GAGAC (SEQ ID NO: 59)

apo-4 = GCGG TGAT GGTG GCAGT (SEQ ID NO: 60)

- 48% perfect homology with Inr

71% match on type of base

(purine vs. pyrimidine)

In Figure 18B, please replace the caption with the following caption:

**FIGURE 18B (SEQ ID NO: 57)**

In Figure 18C, please replace the caption with the following caption:

**FIGURE 18C (SEQ ID NO: 58)**

